

## SEQUENCE LISTING

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&lt;110&gt; Weaver, Lynne

&lt;120&gt; METHODS OF TREATING CHRONIC PAIN USING COMPOSITIONS THAT SPECIFICALLY BIND CD11D (ALPHA-D) INTEGRIN

&lt;130&gt; 27866/39702

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&lt;170&gt; PatentIn version 3.1

&lt;210&gt; 1

&lt;211&gt; 3726

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (3)..(3485)

&lt;223&gt;

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 1130 1135 1140

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Asp Cys Ala Ala Ala Thr Gly Met Cys Gln Pro Ile Pro Leu His Ile  
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Arg Pro Glu Ala Val Asn Met Ser Leu Gly Leu Thr Leu Ala Ala Ser  
 85 90 95

Thr Asn Gly Ser Arg Leu Leu Ala Cys Gly Pro Thr Leu His Arg Val  
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Cys Gly Glu Asn Ser Tyr Ser Lys Gly Ser Cys Leu Leu Leu Gly Ser  
 115 120 125

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Arg Trp Glu Ile Ile Gln Thr Val Pro Asp Ala Thr Pro Glu Cys Pro  
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His Gln Glu Met Asp Ile Val Phe Leu Ile Asp Gly Ser Gly Ser Ile  
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Gly Gln Phe Glu Gly Thr Asp Thr Leu Phe Ala Leu Met Gln Tyr Ser  
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Asn Leu Leu Lys Ile His Phe Thr Phe Thr Gln Phe Arg Thr Ser Pro  
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Ser Gln Gln Ser Leu Val Asp Pro Ile Val Gln Leu Lys Gly Leu Thr  
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Phe Thr Ala Thr Gly Ile Leu Thr Val Val Thr Gln Leu Phe His His  
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Lys Asn Gly Ala Arg Lys Ser Ala Lys Lys Ile Leu Ile Val Ile Thr  
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Asp Gly Gln Lys Tyr Lys Asp Pro Leu Glu Tyr Ser Asp Val Ile Pro  
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Gln Ala Glu Lys Ala Gly Ile Ile Arg Tyr Ala Ile Gly Val Gly His  
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Ala Phe Gln Gly Pro Thr Ala Arg Gln Glu Leu Asn Thr Ile Ser Ser  
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Ala Pro Pro Gln Asp His Val Phe Lys Val Asp Asn Phe Ala Ala Leu  
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Gly Ser Ile Gln Lys Gln Leu Gln Glu Lys Ile Tyr Ala Val Glu Gly  
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Thr Gln Ser Arg Ala Ser Ser Ser Phe Gln His Glu Met Ser Gln Glu  
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Gly Phe Ser Thr Ala Leu Thr Met Asp Gly Leu Phe Leu Gly Ala Val  
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Ser Pro Thr Phe Ile Asn Met Ser Gln Glu Asn Val Asp Met Arg Asp  
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Asn Leu Val Leu Gly Ala Pro Arg Tyr Gln His Thr Gly Lys Ala Val  
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Val Leu Leu Leu Arg Ser Leu Pro Val Leu Lys Val Gly Val Ala Met  
 610 615 620



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Arg Phe Ser Pro Val Glu Val Ala Lys Ala Val Tyr Arg Cys Trp Glu  
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Glu Lys Pro Ser Ala Leu Glu Ala Gly Asp Ala Thr Val Cys Leu Thr  
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Ile Gln Lys Ser Ser Leu Asp Gln Leu Gly Asp Ile Gln Ser Ser Val  
660 665 670

Arg Phe Asp Leu Ala Leu Asp Pro Gly Arg Leu Thr Ser Arg Ala Ile  
675 680 685

Phe Asn Glu Thr Lys Asn Pro Thr Leu Thr Arg Arg Lys Thr Leu Gly  
690 695 700

Leu Gly Ile His Cys Glu Thr Leu Lys Leu Leu Leu Pro Asp Cys Val  
705 710 715 720

Glu Asp Val Val Ser Pro Ile Ile Leu His Leu Asn Phe Ser Leu Val  
725 730 735

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740 745 750

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805 810 815

Ser Leu Tyr Tyr Pro Ala Gly Leu Ser His Arg Arg Val Ser Gly Ala  
820 825 830

Gln Lys Gln Pro His Gln Ser Ala Leu Arg Leu Ala Cys Glu Thr Val  
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Pro Thr Glu Asp Glu Gly Leu Arg Ser Ser Arg Cys Ser Val Asn His  
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Pro Ile Phe His Glu Gly Ser Asn Gly Thr Phe Ile Val Thr Phe Asp  
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Val Ser Tyr Lys Ala Thr Leu Gly Asp Arg Met Leu Met Arg Ala Ser  
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Leu Glu Leu Pro Val Lys Tyr Ala Val Tyr Thr Met Ile Ser Arg Gln  
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Phe Asp Thr Ser Val Tyr Ser Gln Leu Pro Gly Gln Glu Ala Phe  
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Pro Leu Ser  
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